# Spatial Exploration

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May 22, 2018

### **Spatial Exploration**

### Simple Gaussian Process

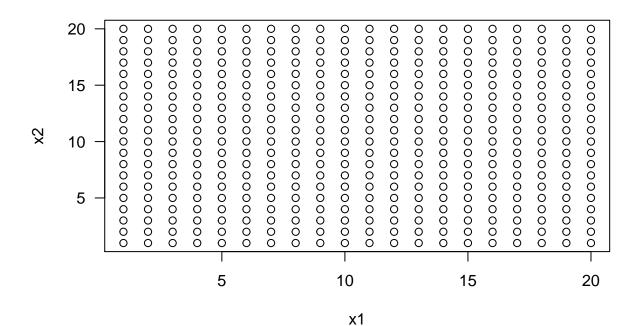
Consider the model:

$$\boldsymbol{y}|\boldsymbol{\theta} \sim MVN(\boldsymbol{X}\boldsymbol{\beta}, \sigma^2\boldsymbol{H}(\phi))$$

where  $H(\phi)$  is the simple exponential case. That is,  $H(\phi) = exp\left(-\frac{d_{ij}}{\phi}\right)$ .

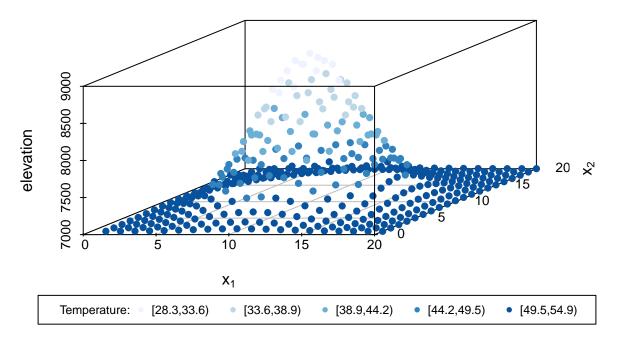
### Generate Spatially Correlated Data

```
set.seed(298032)
grid <- data.frame(expand.grid(1:20, 1:20))
names(grid) <- c("x1", "x2")
plot(grid, las = 1)</pre>
```



```
sp.grid <- grid</pre>
coordinates(sp.grid) <- ~x1 + x2</pre>
d <- rgeos::gDistance(sp.grid, byid = T)</pre>
# create spatial covariate - elevation
grid$elevation <- with(grid, 2000 * \exp(-(1/20) * ((x1 - 10)^2 + (x2 - 10)^2)) +
    7000)
# create response - temperature #b0 = 140, b1 = -1/80, sigma2 = 1, phi = 10
temp <- with(grid, rmvnorm(1, mean = 140 - (1/80) * elevation, sigma = 1 * exp(-d/10))
# final spatial df
garnet.df <- data.frame(cbind(grid, temp = as.numeric(temp)))</pre>
MyPalette <- brewer.pal(5, "Blues")</pre>
palette(MyPalette)
split.obs <- cut(garnet.df$temp, 5, right = F)</pre>
cuts <- levels(split.obs)</pre>
par(mar = c(10, 4, 4, 2) + 0.1)
scatterplot3d::scatterplot3d(x = grid$x1, y = grid$x2, z = grid$elevation, pch = 16,
    color = MyPalette[as.numeric(split.obs)], main = "Garnet Mountain", xlab = expression("x"[1]),
    ylab = expression("x"[2]), zlab = "elevation", bg = "red")
legend("bottom", legend = c("Temperature:", cuts), pch = c(NA, rep(16, 5)),
    col = c("white", MyPalette), inset = -0.35, xpd = T, horiz = T, cex = 0.75)
```

### **Garnet Mountain**



```
# check it out plot3d(x = garnet.df$x1, y = garnet.df$x2, z = garnet.df$elevation, main = Garnet Mountain, col = garnet Mountain
```

```
# MyPalette[as.numeric(split.obs)], size = 5, xlab = '', ylab = '', zlab =
# '') rgl.bbox(xlen = 0, ylen = 0, color = c('orange')) legend3d('topright',
# legend = cuts, pch = 16, col = MyPalette, cex=.7, inset=c(0.05))
# snapshot3d(filename = '3dplot.png', fmt = 'png')

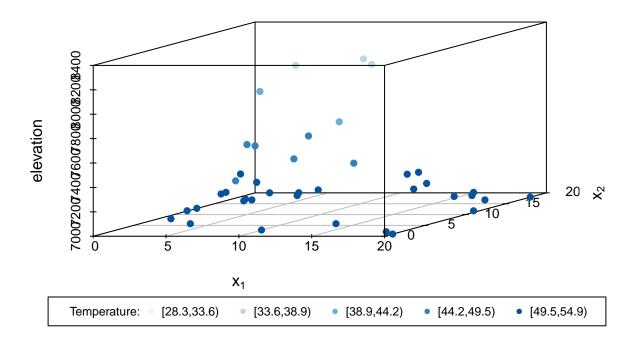
# sample points
set.seed(16240)
id <- sample(400, 40)

samp <- garnet.df[sort(id), ]
split.samp.obs <- cut(samp$temp, c(28.3, 33.6, 38.9, 44.2, 49.5, 54.9), right = F)

scatterplot3d::scatterplot3d(x = samp$x1, y = samp$x2, z = samp$elevation, pch = 16, color = MyPalette[as.numeric(split.samp.obs)], main = "Sample of Garnet Mountain", xlab = expression("x"[1]), ylab = expression("x"[2]), zlab = "elevation", bg = "red")

legend("bottom", legend = c("Temperature:", cuts), pch = c(NA, rep(16, 5)), col = c("white", MyPalette), inset = -0.35, xpd = T, horiz = T, cex = 0.75)</pre>
```

# **Sample of Garnet Mountain**



### Sampler

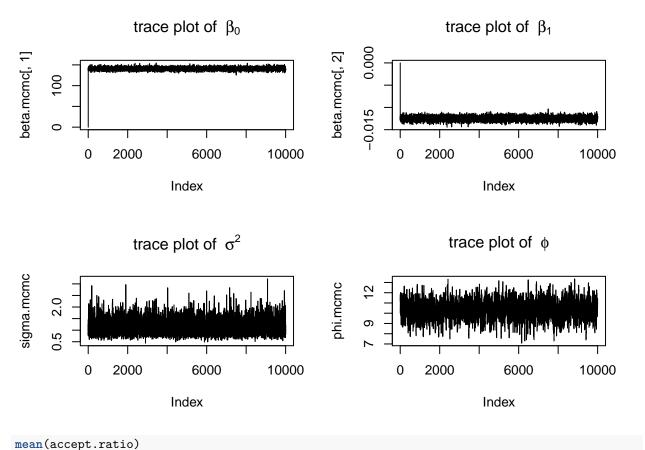
Priors:

```
p(\beta) = \text{mvnormal}(\boldsymbol{\mu}_0, \boldsymbol{\Sigma}_0)p(\sigma^2) = \text{inverse-gamma}(\frac{\nu_0}{2}, \frac{\sigma_0^2}{2})p(\phi) = \text{gamma}(a, b)
```

Note: Ended up placing a normal (10,1) prior on  $\phi$ ... very informative, couldn't get it to behave otherwise.

```
# priors
n <- 40
mu0 \leftarrow rep(0, 2)
Sigma0 <- 1000 * diag(2)
nu0 <- 1e-04
sigma20 <- 10
# nu0 <- 6 sigma20 <- 4
a < -1e-04
b < -1e-04
Sigma0.inv <- solve(Sigma0)</pre>
sp.samp.grid \leftarrow data.frame(x1 = samp$x1, x2 = samp$x2)
coordinates(sp.samp.grid) <- ~x1 + x2</pre>
dist.mat <- rgeos::gDistance(sp.samp.grid, byid = T)</pre>
y <- samp$temp
X <- model.matrix(~elevation, data = samp)</pre>
# setup sampler
num.mcmc <- 10000
step.size <- 0.5
beta.mcmc <- matrix(0, num.mcmc, 2)</pre>
sigma.mcmc <- rep(0, num.mcmc)</pre>
phi.mcmc <- rep(0, num.mcmc)</pre>
accept.ratio <- rep(0, num.mcmc)</pre>
# initialize sampler
sigma.mcmc[1] <- 1
phi.mcmc[1] <- 10
for (i in 2:num.mcmc) {
    # gibbs step for beta and sigma
    # sample beta | sigma, phi
    Sigma <- sigma.mcmc[i - 1] * exp(-dist.mat/phi.mcmc[i - 1])</pre>
    Sigma.inv <- solve(Sigma)</pre>
    A <- solve(t(X) %*% Sigma.inv %*% X + Sigma0.inv)
    B <- t(X) %*% Sigma.inv %*% y - Sigma0.inv %*% mu0
    beta.mcmc[i, ] <- mvtnorm::rmvnorm(1, mean = A %*% B, sigma = A)
    # sample sigma | beta, phi
    resid <- y - X %*% beta.mcmc[i, ]</pre>
    H <- exp(-dist.mat/phi.mcmc[i - 1])</pre>
```

```
H.inv <- solve(H)</pre>
    SSR <- t(resid) %*% H.inv %*% resid
    sigma.mcmc[i] <- LearnBayes::rigamma(1, (nu0 + n)/2, (SSR + sigma20)/2)
    # metropolis step for phi
    phi.s <- phi.mcmc[i - 1]</pre>
    phi.star <- phi.mcmc[i - 1] + rnorm(1, mean = 0, sd = sqrt(step.size))</pre>
    num <- mvtnorm::dmvnorm(y, mean = X %*% beta.mcmc[i, ], sigma = sigma.mcmc[i] *</pre>
        exp(-dist.mat/phi.star), log = T) + dnorm(phi.star, mean = 10, sd = 1,
        log = T)
    \# dunif(phi.star, 0, max(dist.mat), log = T) \#dgamma(phi.star, a, 1 / b, log
    \# = T) \#dnorm(phi.star, mean = 10, sd = 1, loq = T) \#dexp(phi.star, 1/a, loq = T)
    \# = T
    denom <- mvtnorm::dmvnorm(y, mean = X %*% beta.mcmc[i, ], sigma = sigma.mcmc[i] *</pre>
        exp(-dist.mat/phi.s), log = T) + dnorm(phi.s, mean = 10, sd = 1, log = T)
    # dunif(phi.s, 0, max(dist.mat), log = T) # dqamma(phi.s, a, 1 / b, log = T)
    ##dnorm(phi.s, mean = 10, sd = 1, loq = T) #dexp(phi.s, 1/a, loq = T)
    log.r <- num - denom
    if (log(runif(1)) < log.r) {</pre>
        phi.mcmc[i] <- phi.star</pre>
        accept.ratio[i] <- 1</pre>
    } else {
        phi.mcmc[i] <- phi.s</pre>
    if (i\%500 == 0) {
        message("Progress: ", bquote(.(i)), "th iteration of chain")
        message(bquote(.(round(i/(num.mcmc), 5) * 100)), "% through chain")
    }
}
par(mfrow = c(2, 2))
plot(beta.mcmc[, 1], type = "l", main = expression("trace plot of " ~ beta[0]))
plot(beta.mcmc[, 2], type = "l", main = expression("trace plot of " ~ beta[1]))
plot(sigma.mcmc, type = "l", main = expression("trace plot of " ~ sigma^2))
plot(phi.mcmc, type = "l", main = expression("trace plot of " ~ phi))
```



moun (accept:ratio)

## [1] 0.7658

#### Posterior Predictive Checks

To predict across a vector of unobserved locations,  $S_0 = \{s_{01}, s_{02}, ..., s_{0m}\}$ , we have:

$$p(\boldsymbol{y}_0|\boldsymbol{y},\boldsymbol{X},\boldsymbol{X}_0) = \int p(\boldsymbol{y}_0|\boldsymbol{y},\boldsymbol{\theta},\boldsymbol{X}_0) p(\boldsymbol{\theta}|\boldsymbol{y},\boldsymbol{X}) d\boldsymbol{\theta} \approx \frac{1}{G} \sum_{g=1}^G p(\boldsymbol{y}_0|\boldsymbol{y},\boldsymbol{\theta}^{(g)},\boldsymbol{X}_0)$$

```
# predict across population grid
X0 <- model.matrix(~elevation, data = garnet.df)

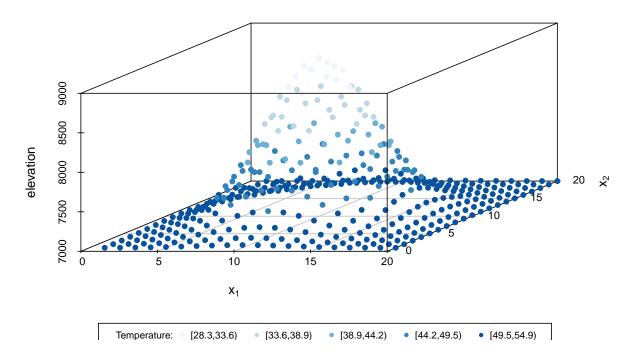
# posterior predictive distribution
pred <- matrix(0, num.mcmc, nrow(garnet.df))
for (i in 1:dim(beta.mcmc)[1]) {
    mean <- X0 %*% beta.mcmc[i, ]
    H <- exp(-d/phi.mcmc[i])
    var <- sigma.mcmc[i] * H

    pred[i, ] <- rmvnorm(1, mean = mean, sigma = var)

if (i%%500 == 0) {
    message("Progress: ", bquote(.(i)), "th iteration")</pre>
```

```
message(bquote(.(round(i/(num.mcmc), 5) * 100)), "% through")
    }
}
pred.burn <- pred[500:dim(pred)[1], ]</pre>
pred.temp <- colMeans(pred.burn)</pre>
pred.df <- cbind(garnet.df[, 1:3], pred.temp)</pre>
split.pred.obs \leftarrow cut(pred.df_pred.temp, c(28.3, 33.6, 38.9, 44.2, 49.5, 54.9),
    right = F)
par(mfrow = c(2, 1))
scatterplot3d::scatterplot3d(x = pred.df$x1, y = pred.df$x2, z = pred.df$elevation,
    pch = 16, color = MyPalette[as.numeric(split.pred.obs)], main = "Predicted Garnet Mountain",
    xlab = expression("x"[1]), ylab = expression("x"[2]), zlab = "elevation",
    bg = "red")
legend("bottom", legend = c("Temperature:", cuts), pch = c(NA, rep(16, 5)),
    col = c("white", MyPalette), inset = -0.35, xpd = T, horiz = T, cex = 0.75)
scatterplot3d::scatterplot3d(x = grid$x1, y = grid$x2, z = grid$elevation, pch = 16,
    color = MyPalette[as.numeric(split.obs)], main = "Garnet Mountain", xlab = expression("x"[1]),
    ylab = expression("x"[2]), zlab = "elevation", bg = "red")
legend("bottom", legend = c("Temperature:", cuts), pch = c(NA, rep(16, 5)),
   col = c("white", MyPalette), inset = -0.35, xpd = T, horiz = T, cex = 0.75)
```

### **Predicted Garnet Mountain**



# **Garnet Mountain**

